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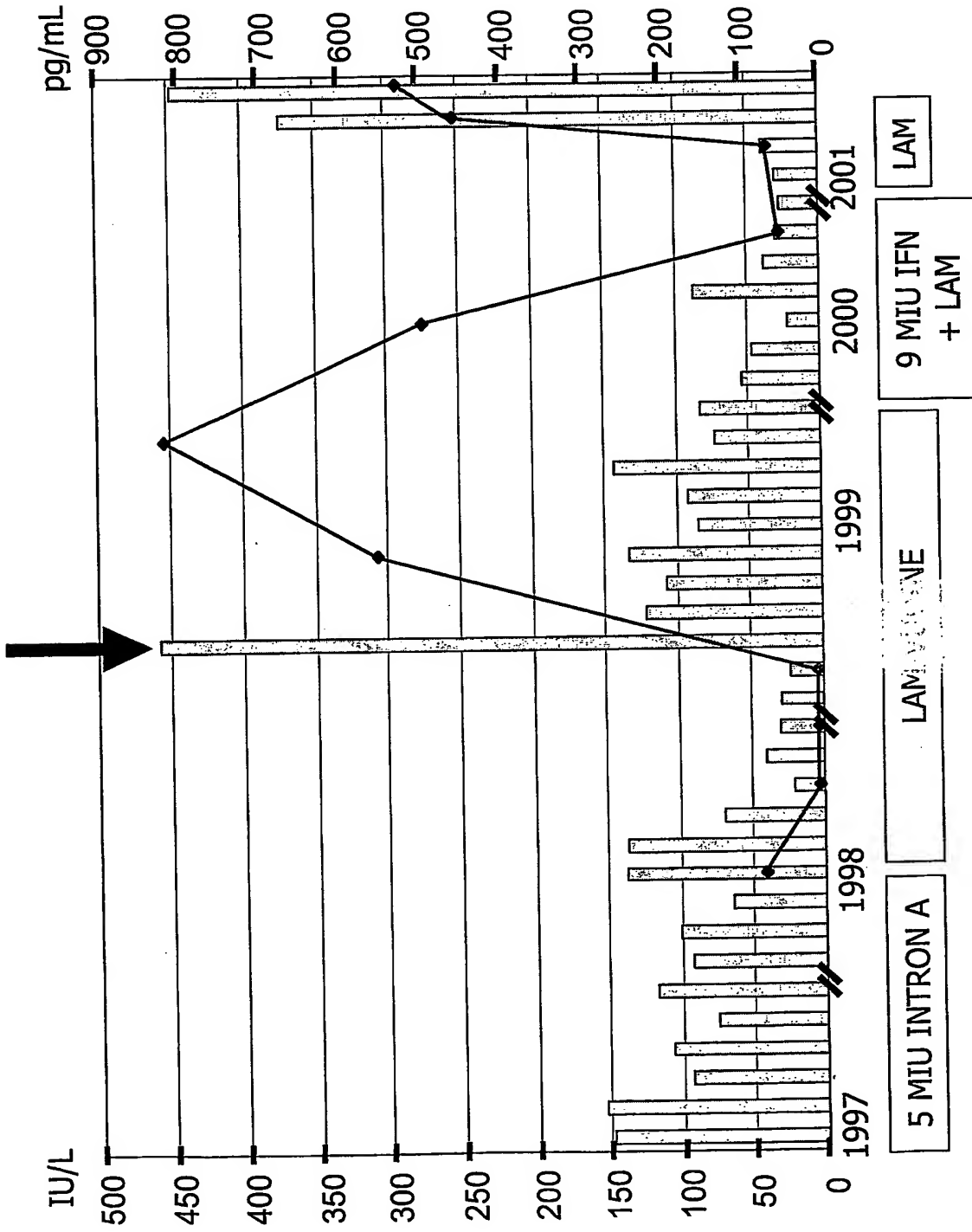


FIGURE 1

CONFIRMATION COPY

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RT		100		110		120						
HBVD	RT	AMP	HLL	VGSS	GLS	RYVAR	LSS	NSR	IF	NYQ		
P7RT											
RT		130		140		150						
HBVD	RT	HGT	MQN	LHD	SCSR	NLY	VSL	LLLY	QTF	GRK		
P7RT											
RT		160		170		180						
HBVD	RT	LHL	YSH	PIIL	GFR	KIP	MGV	GLS	PFL	L	AQF	
P7RT										M	...
RT		190		200		210						
HBVD	RT	TSA	ICS	VVRR	AF	PE	GLA	FSY	M	DDV	VLGAK	
P7RT										S
RT		220		230		240						
HBVD	RT	SVQ	HLES	LFT	AVT	NFL	LSL	GIH	LNP	NKTK		
P7RT											
RT		250		256								
HBVD	RT	RWG	YSL	HFM	GYV	IGC						
P7RT											

HBVD RT = SEQ ID NO:1

P7RT = SEQ ID NO:4

FIGURE 2

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S 90 100 110
 | | |
HBVD S LCLIFLLVLLDYQGMLPVCPLIPGSSTTS
P7S

 120 130 140
 | | |
HBVD S TGPCRTCTTPAQGTSMYPSCCCTKPSDGN
P7S

 150 160 170
 | | |
HBVD S CTCIPIPSWAFGKFLWEWASARFSWLSL
P7S

 180 190 200
 | | |
HBVD S LVEFVQWVGLSPTVWLSVIWMMWYWGPS
P7S V.....

 210 220 226
 | | |
HBVD S LYSILSPFLPLLPIFFCLWVYI
P7S

HBVD S = SEQ ID NO:2

P7S = SEQ ID NO:5

FIGURE 3

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HBVD 420 gctatgcctcatcttcttggttggttcttctggactatcaa
P7

HBVD 460 ggtatggtgcccgtttgtcctctaattccaggatcttcaa
P7

HBVD 500 ctaccagcacgggaccatgcagaacctgcacgactcctgc
P7

HBVD 540 tcaaggaacctctatgtatccctcctggttgctgtaccaa
P7

HBVD 580 ccttcggacggaaattgcacctgtattcccatcccatcat
P7

HBVD 620 cctgggctttcggaaaattcctatgggagtgggcctcagc
P7

HBVD 660 ccgtttctc **c** tggctcagtttactagtgccatttgttcag
P7 **a**

HBVD 700 tggttcgtagggctttcccccactgtttggctttcagtt**a**
P7

HBVD 740 ta **t g** gatgat gtggtattgggggccaagtctgtacagcat
P7 .. **g t**

HBVD 780 cttgagtccttttttaccgctgttaccaattttctt **c** tgt
P7 **t**

HBVD 820 ctttgggtatacatttaaaccctaacaaaaacaaaaagatg
P7

HBVD 860 gggttactctttacatttcattggg **c** tatgtcattggatgt
P7 **g**

HBVD = SEQ ID NO:3

P7 = SEQ ID NO:6

FIGURE 4